

Visual Alignment

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Intro to Visual Alignment Program (visualalign)

- Purpose
- Event reconstruction
- Track selection
- Alignment procedure
- Displaying residuals at step
- Alignment step options

Purpose

- Visualize alignment problems
 - 3D residuals
 - Display selected residuals
- Develop alignment strategies graphically
 - Selection of alignment objects / parameters
 - Selection of tracks to use
 - Specification of sequence of such alignments
- Record strategy for reuse and modification
- Align all DØ detectors

Event Reconstruction

- Load Mutable Offline Geometry
- Must redo Reco after each alignment step
- Local clusters do not need redo
- Repeatedly run D0Reco with I,O files?
- Save stripped down events in memory?
- Call D0Reco from alignment program?

Track Selection

- Tracks with $p_t > \text{say } 5 \text{ GeV}/c$ for barrel
- Tracks with high p_z for disks (minbias?)
- Might use several samples at once.
- Do we need special triggers?
- Multiple tracks vertexed to interaction?

Alignment Procedure

- Reconstruct events
- Select objects / geometry transforms
- Select components of transforms
- Select track sample(s)
- Display residuals
- Try step, accept or reject it
- Repeat above until alignment done

Displaying Residuals at Step

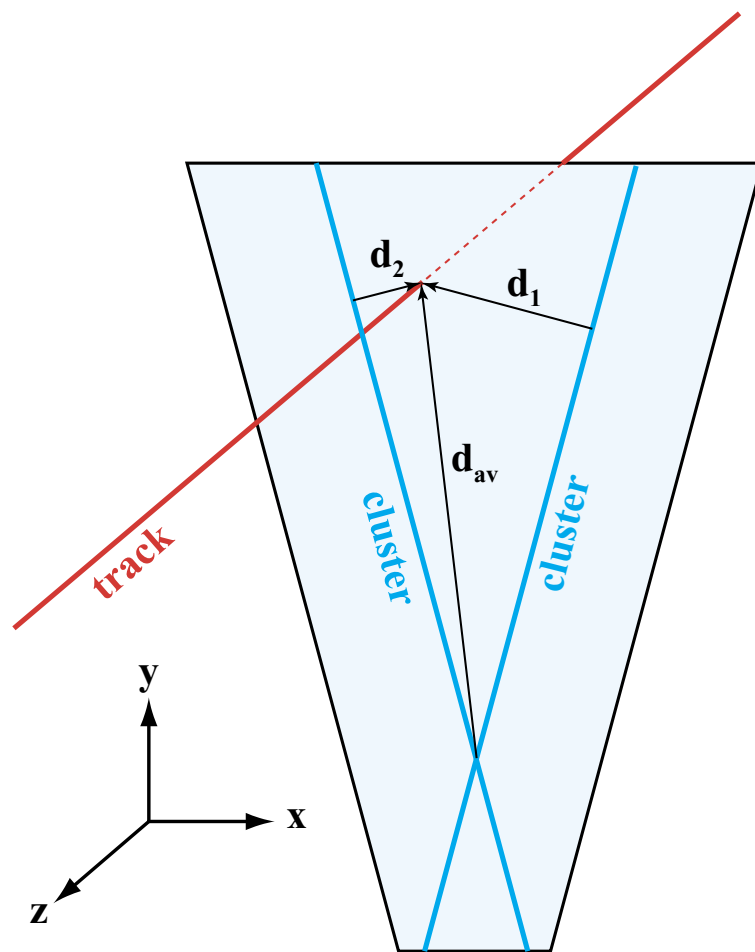
- Display 3D vectors (with error ellipse?)
- Cut on significance, e.g. 3 sigma from zero
- At each alignment step:
 - Display selected objects / transforms
 - Display selected track sample(s)
 - Display related residual field
 - Display χ^2 , Rot, Trans for each transform

Alignment Step Options

- Try this alignment step
 - Rot, Trans to obtain minimum χ^2
 - Display Residuals after this step
 - Either: Reject this step
 - Back up to selections for this step
 - Or: Accept this step
 - Log selections for this step
 - Begin selections for next step
- Manually move object; view χ^2 , Rot, Trans

Progress on visualalign

- 3D residual concept tested
- Some Classes written, in CVS
 - Residual
 - SMTResidual (prototype XXXResidual)
 - ResidualField written
- Trying to read events
- Immediate goal: Display some residuals
- Longer term goal: Align 10% test



The $\chi^2(\mathbf{Q}, \mathbf{h})$ that \mathbf{d}'_i , the residuals after rotation \mathbf{R} ($\mathbf{Q} = \mathbf{R} - \mathbf{1}$) and translation \mathbf{h} , are equal to zero within their error matrices \mathbf{E}_i is

$$\begin{aligned}\chi^2(\mathbf{Q}, \mathbf{h}) &= \sum_i \mathbf{d}'_i{}^\dagger \mathbf{E}_i^{-1} \mathbf{d}'_i, \\ \mathbf{d}'_i &= \mathbf{d}_i - \mathbf{Q} \mathbf{s}_i - \mathbf{h}, \\ \mathbf{Q} &= \mathbf{R} - \mathbf{1}.\end{aligned}$$

Differentiating χ^2 with respect to the components of \mathbf{Q} and \mathbf{h} and setting the differentials equal to zero yields six equations in the components of \mathbf{Q} and \mathbf{h} :

$$\begin{aligned}\sum_{m=1}^3 a_{jm} \theta_m + \sum_{n=1}^3 b_{jn} h_n &= c_j, \quad j=1,3, \\ \sum_{m=1}^3 d_{jm} \theta_m + \sum_{n=1}^3 e_{jn} h_n &= f_j, \quad j=1,3,\end{aligned}$$

$$a_{jm} = \sum_i \mathbf{s}_i^\dagger \mathbf{Q}_j^\dagger \mathbf{E}_i^{-1} \mathbf{Q}_m \mathbf{s}_i, \quad b_{jn} = \sum_i \mathbf{s}_i^\dagger \mathbf{Q}_j^\dagger \mathbf{E}_i^{-1} \mathbf{u}_n, \quad c_j = \sum_i \mathbf{s}_i^\dagger \mathbf{Q}_j^\dagger \mathbf{E}_i^{-1} \mathbf{d}_i,$$

$$d_{jm} = \sum_i \mathbf{u}_j^\dagger \mathbf{E}_i^{-1} \mathbf{Q}_m \mathbf{s}_i, \quad e_{jn} = \sum_i \mathbf{u}_j^\dagger \mathbf{E}_i^{-1} \mathbf{u}_n, \quad f_j = \sum_i \mathbf{u}_j^\dagger \mathbf{E}_i^{-1} \mathbf{d}_i.$$

Assuming no rotation, the translation vector \mathbf{h} which minimizes χ^2 is

$$\mathbf{h} = \left(\sum_i \mathbf{E}_i^{-1} \right)^{-1} \sum_i \mathbf{E}_i^{-1} \mathbf{d}_i,$$

which is the weighted average of the residuals, where the weighting includes the effects of correlations.

\mathbf{d}_i is residual
 \mathbf{t}_i is track
 \mathbf{s}_i is cluster

$$\begin{aligned}\mathbf{d}_i &= \mathbf{t}_i - \mathbf{s}_i \\ \mathbf{s}'_i &= \mathbf{R} \mathbf{s}_i + \mathbf{h} \\ \mathbf{d}'_i &= \mathbf{t}_i - \mathbf{s}'_i \\ &= \mathbf{d}_i + \mathbf{s}_i - \mathbf{s}'_i \\ &= \mathbf{d}_i + \mathbf{s}_i - \mathbf{R} \mathbf{s}_i - \mathbf{h} \\ &= \mathbf{d}_i - (\mathbf{R} - \mathbf{1}) \mathbf{s}_i - \mathbf{h} \\ &= \mathbf{d}_i - \mathbf{Q} \mathbf{s}_i - \mathbf{h}\end{aligned}$$